

APPROVED	O.G. FIG.
BY	CLASS      SUBCLASS
DRAFTSMAN	

08/973303

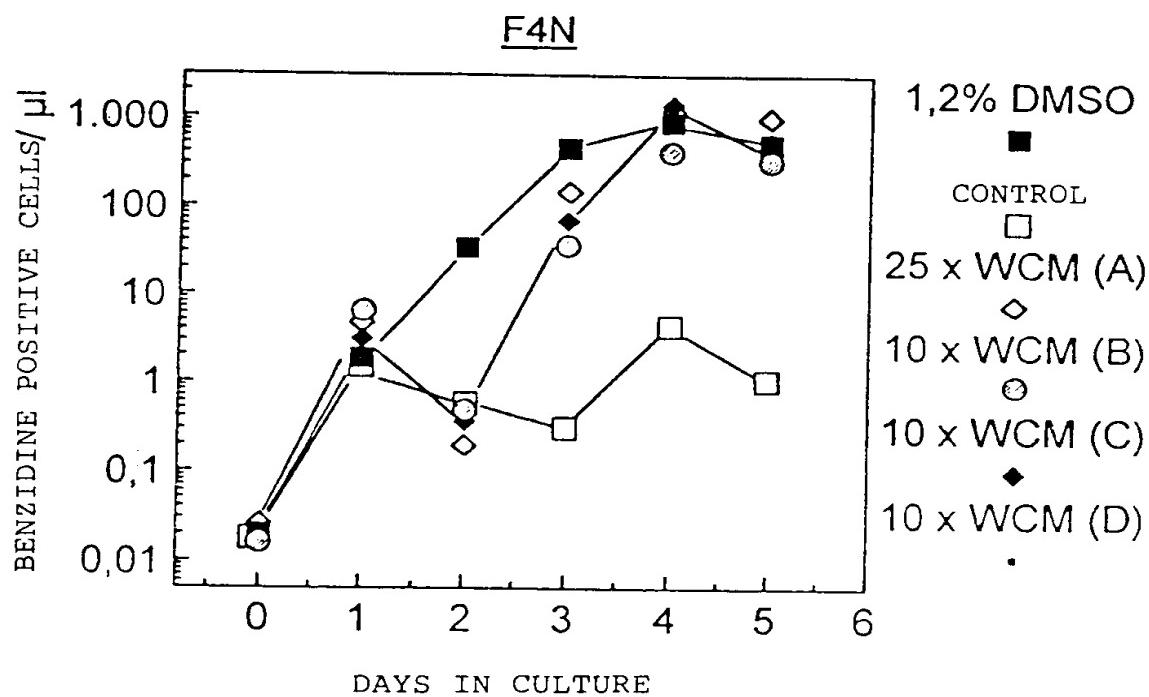


FIGURE 1

APPROVED BY	O.G. FIG.
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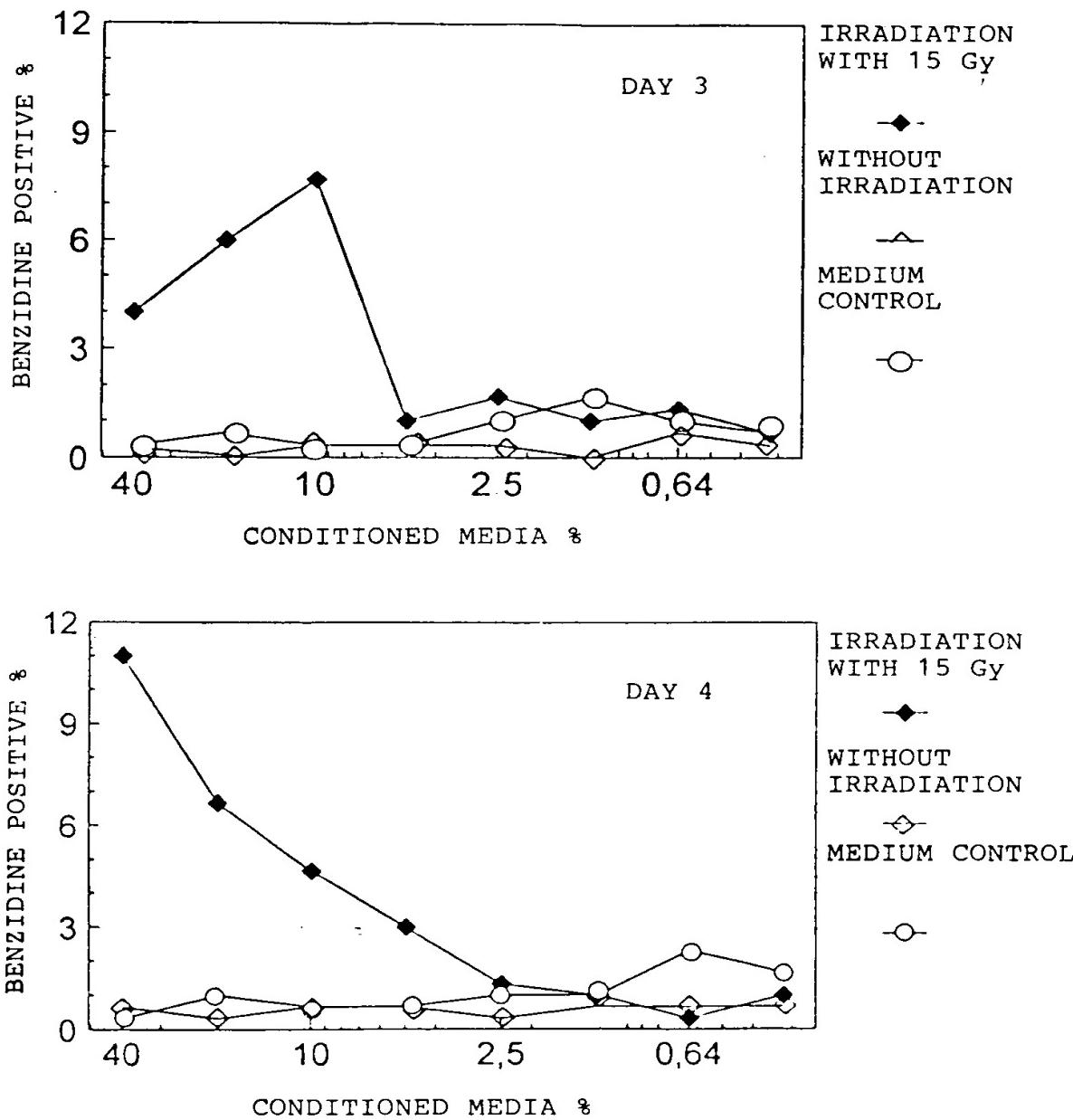


FIGURE 2

APPROVED BY	O.G. FIG.
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K 562

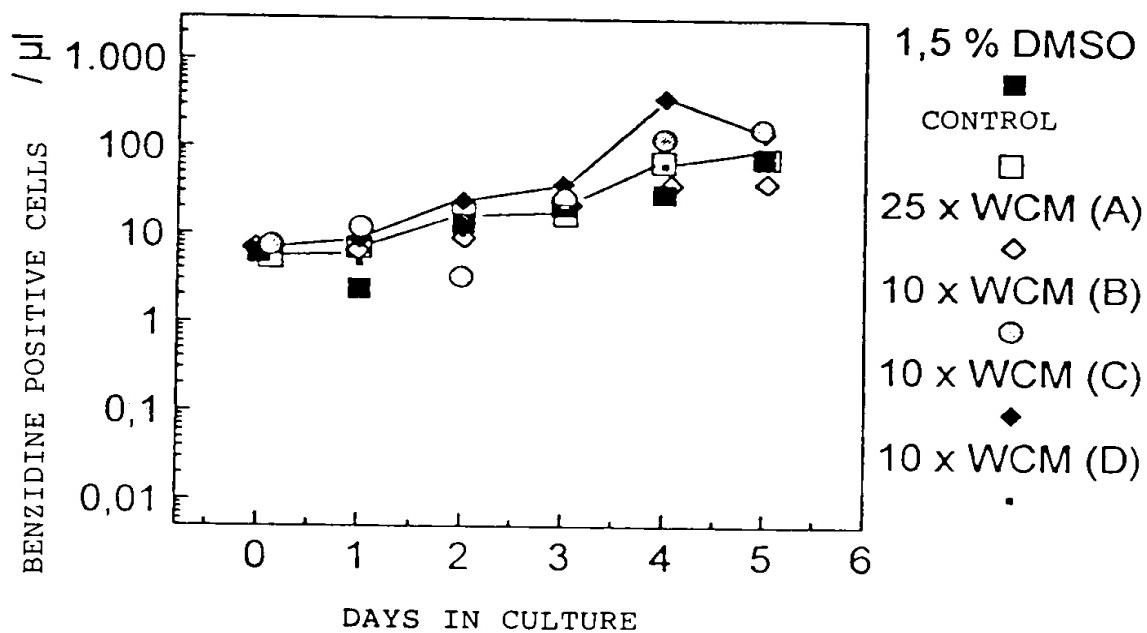


FIGURE 3

APPROVED BY	O.G. FIG.	
DRAFTSMAN	CLASS	SUBCLASS

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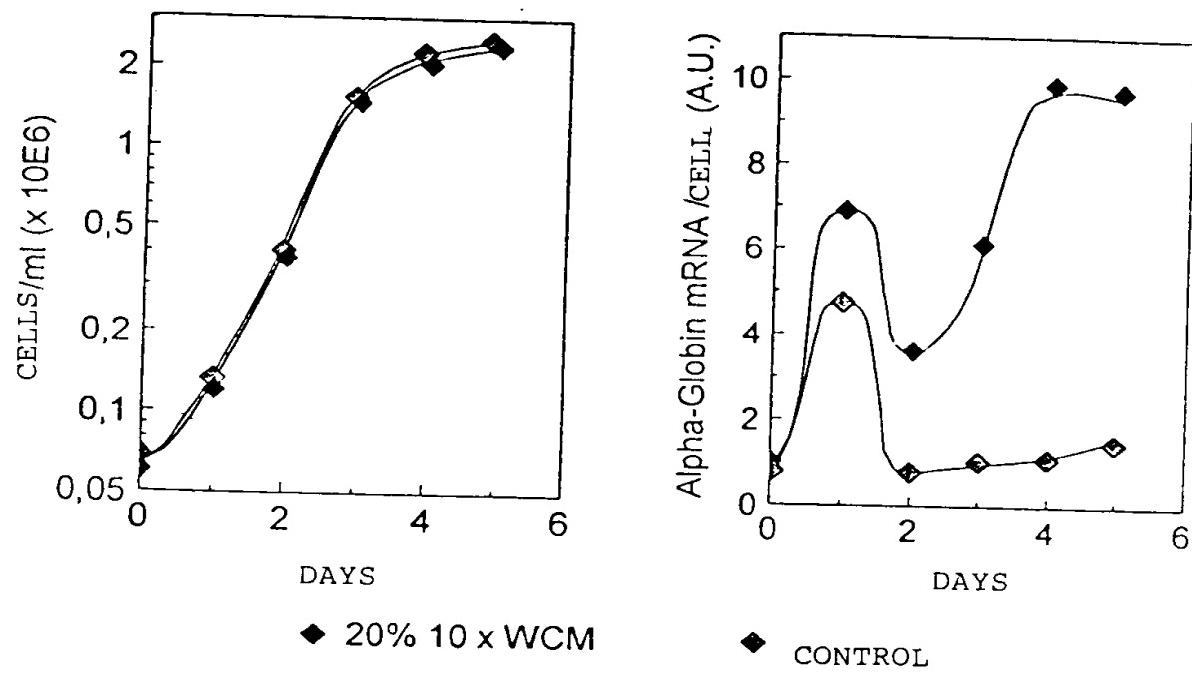


FIGURE 4

APPROVED BY	O.G. FIG.
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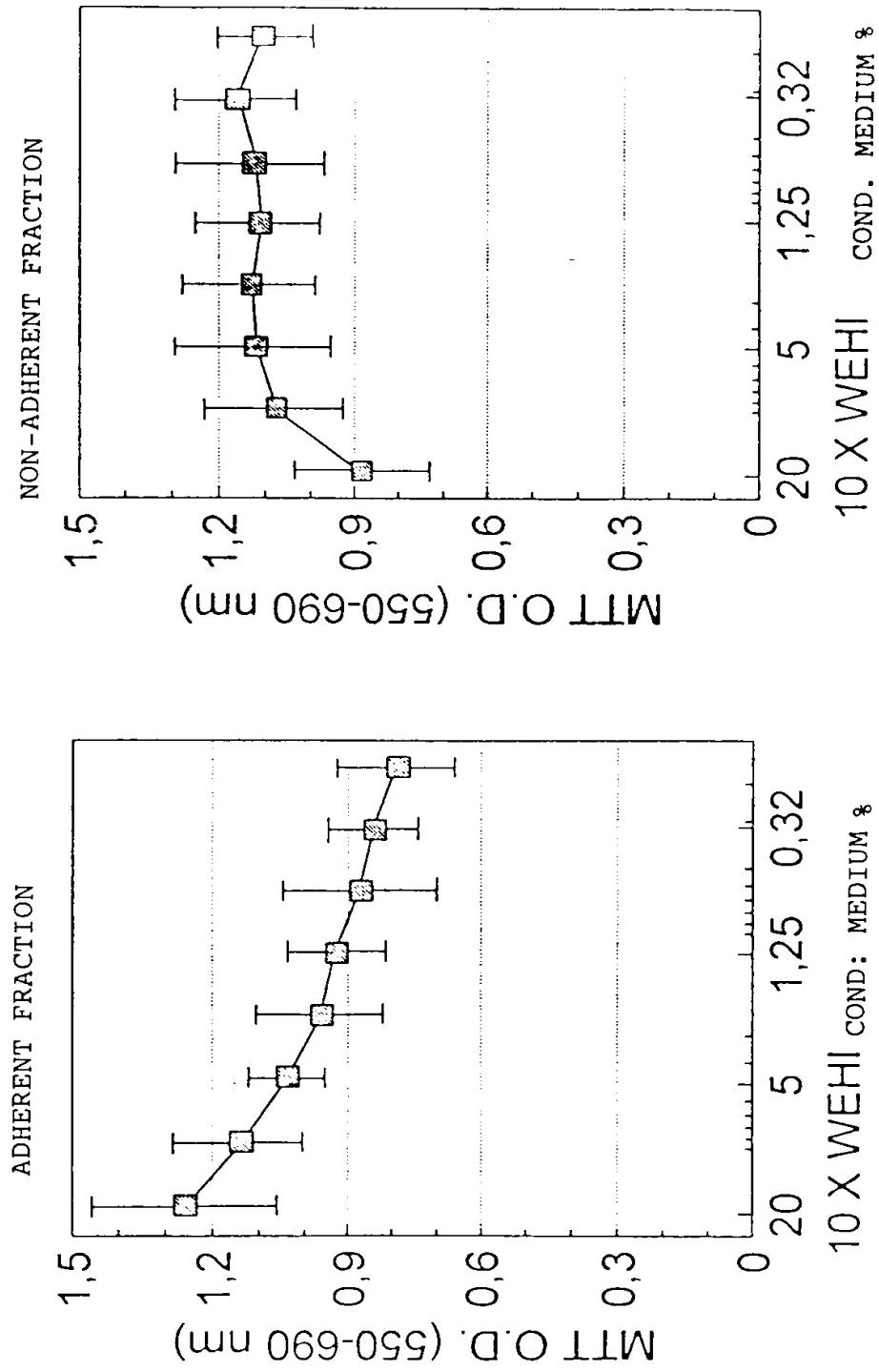


FIGURE 5

APPROVED	O.G. FIG.
BY	CLASS
DRAFTSMAN	SUBCLASS

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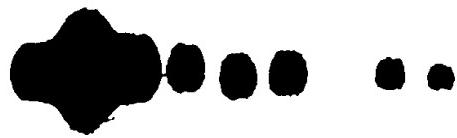
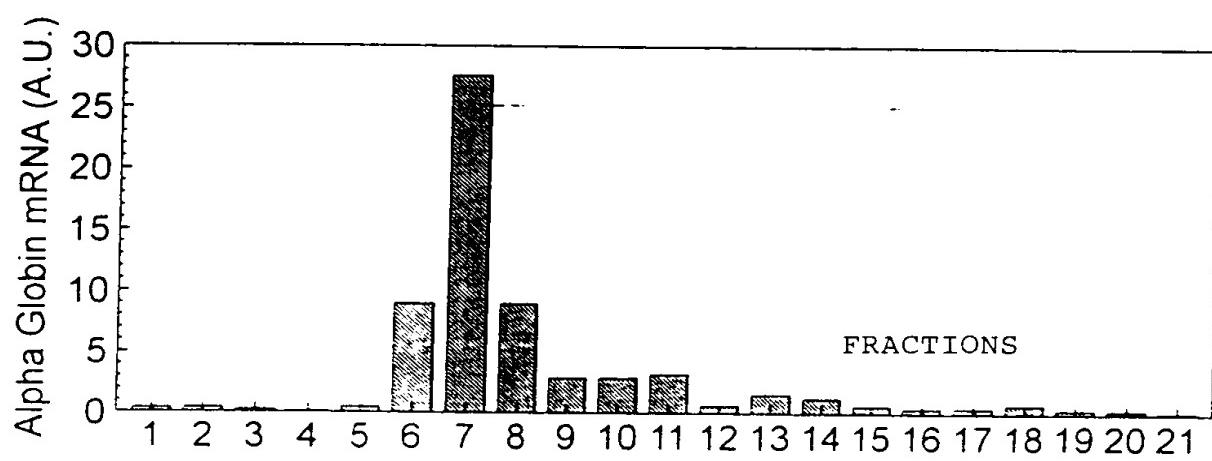
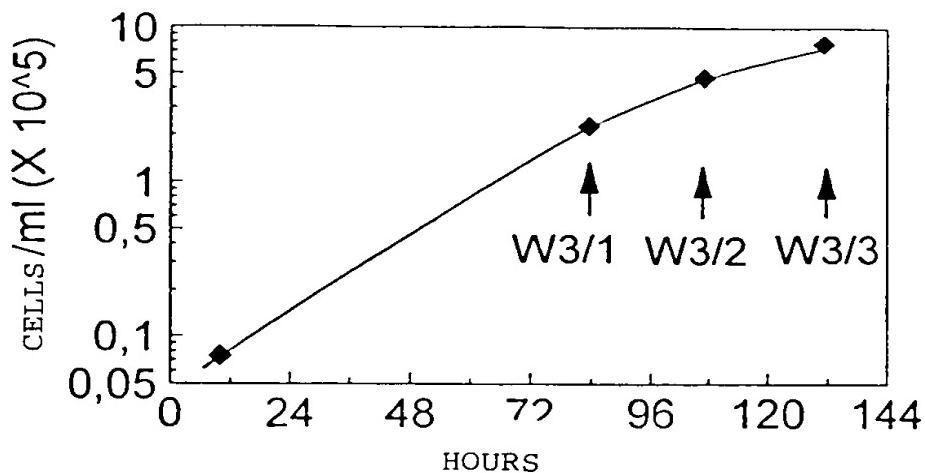


FIGURE 6

APPROVED	O.G. FIG.
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CELL DENSITY



INDUCTION FOR DIFFERENTIATION

(B 8/3, DAY 3)

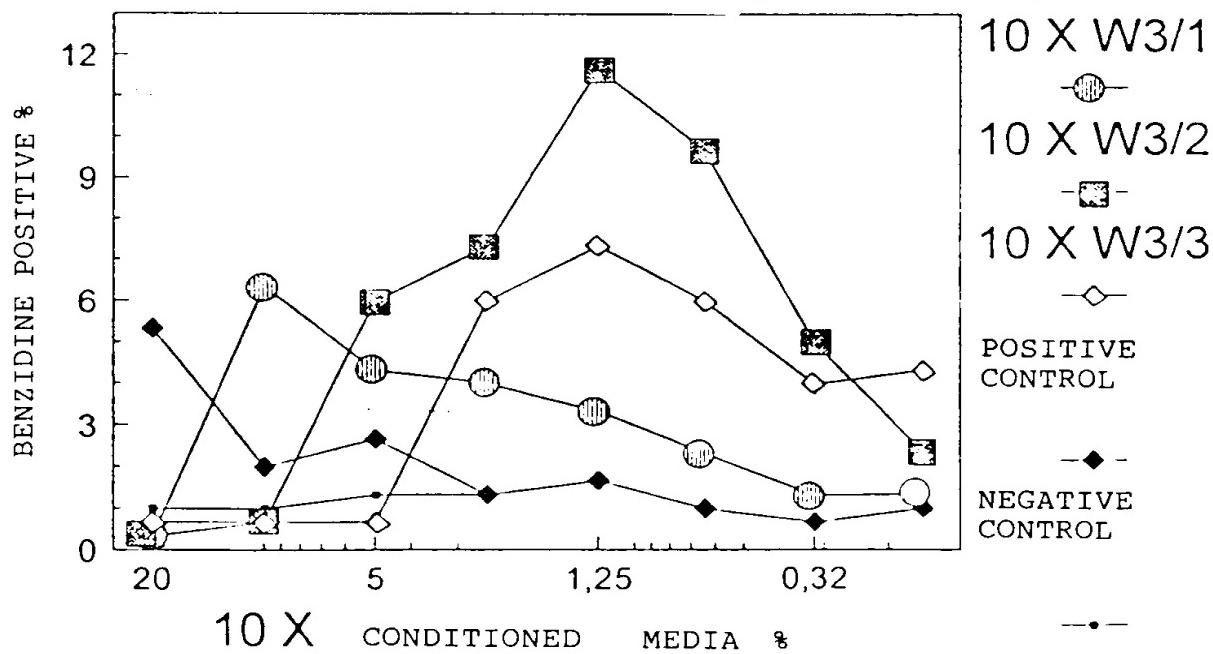


FIGURE 7

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APPROVED BY	O.G. FIG.
DRAFTSMAN	CLASS SUBCLASS

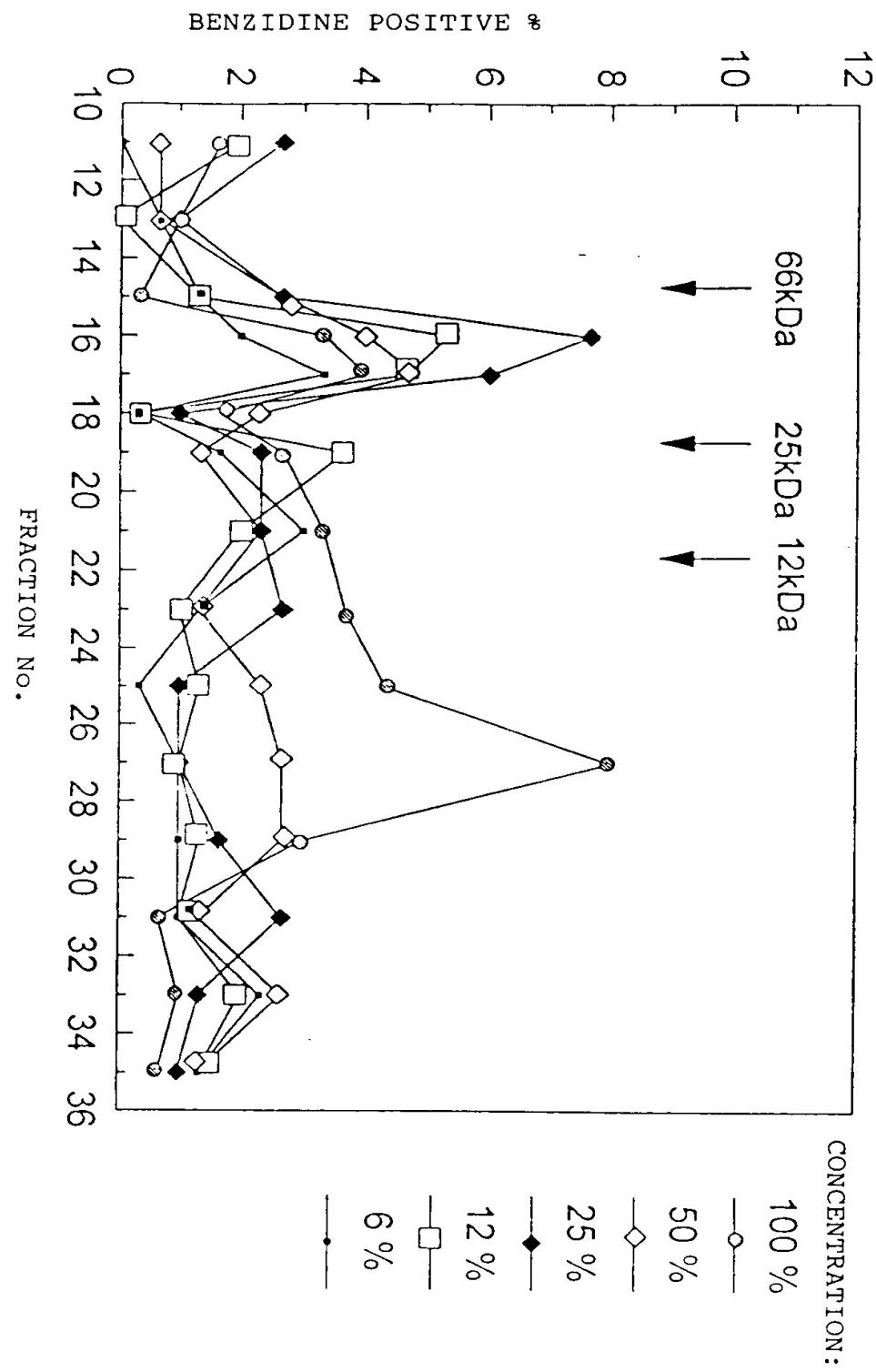


FIGURE 8

APPROVED BY	O.G. FIG.
DRAFTSMAN	CLASS SUBCLASS

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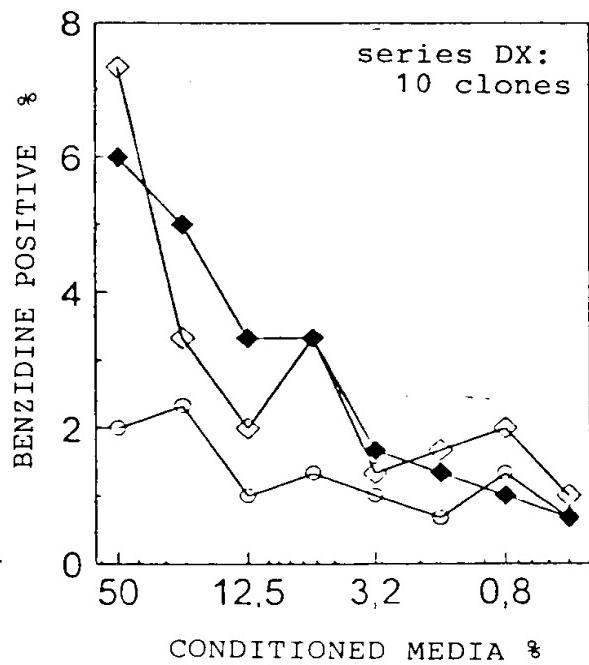
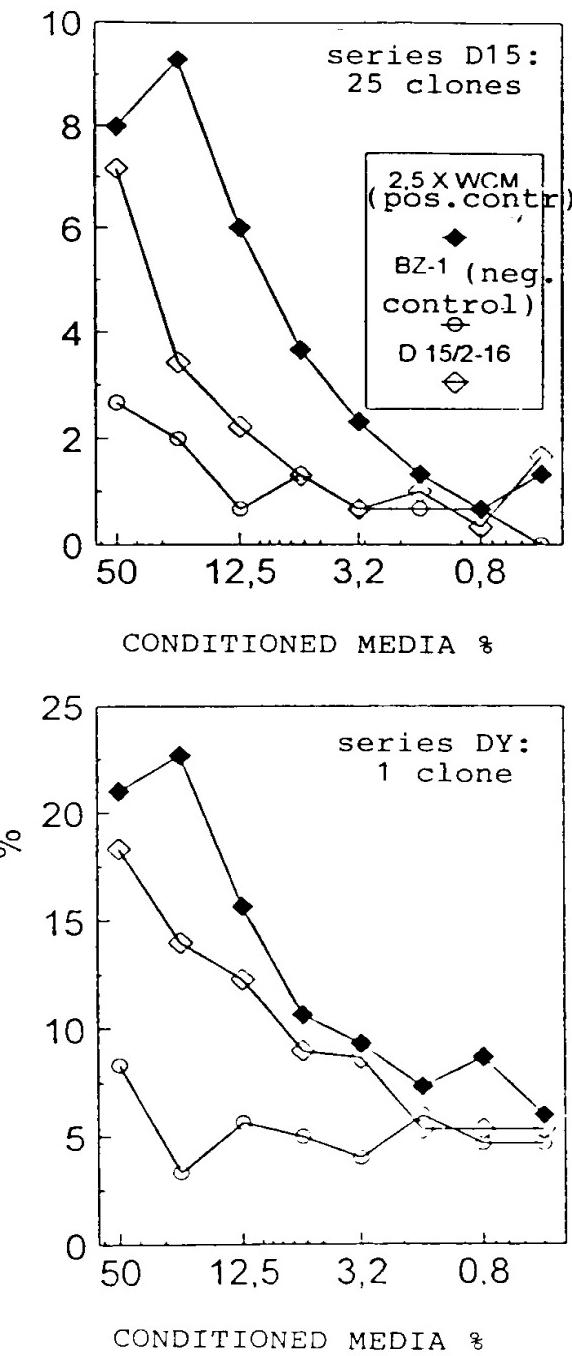
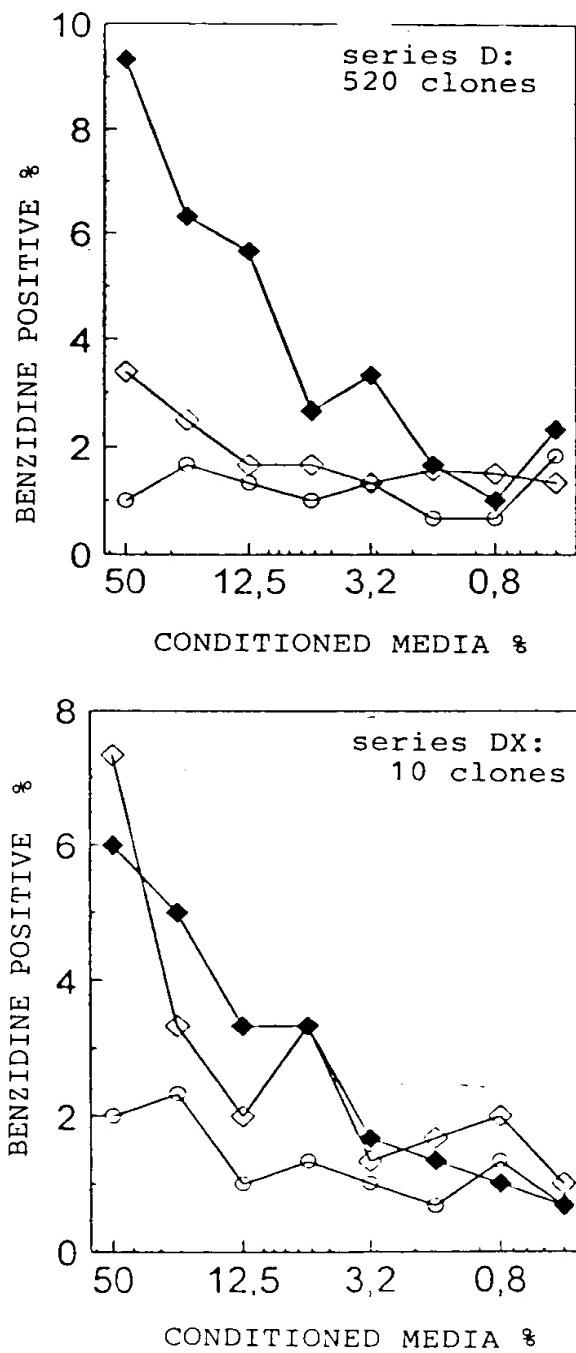


FIGURE 9

APPROVED BY	O.G. FIG.
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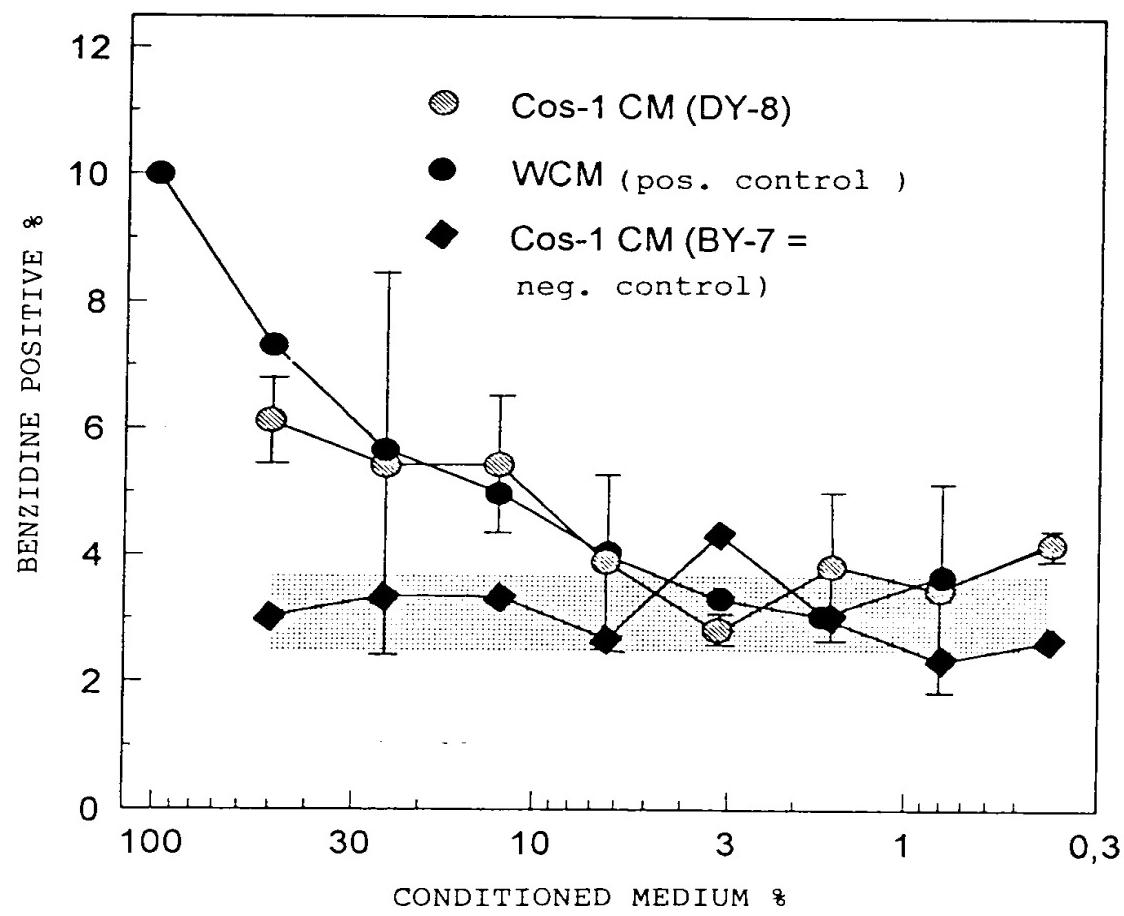


FIGURE 10

APPROVED BY	O.G. FIG.
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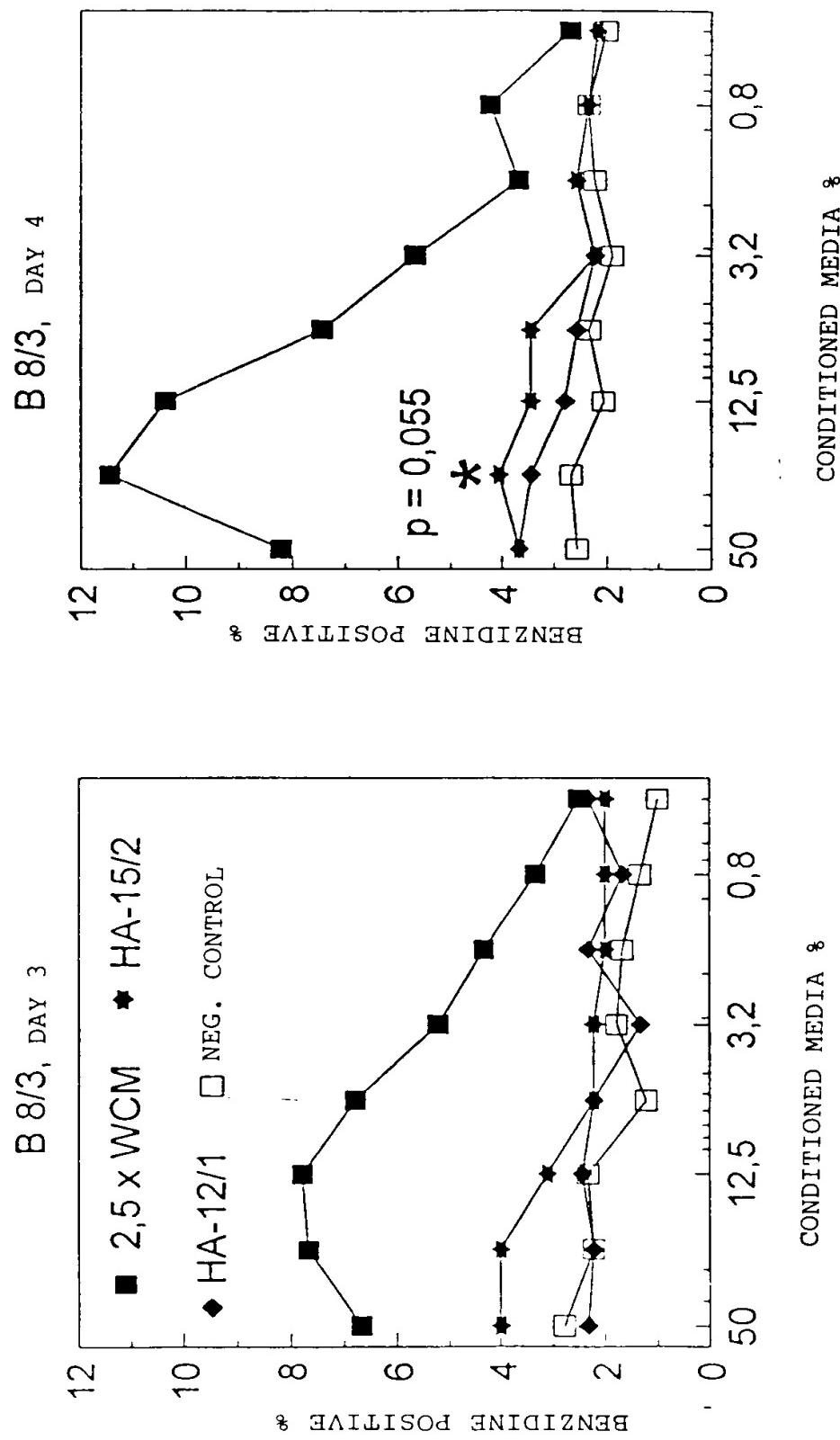


FIGURE 11

APPROVED BY	O.G. FIG.
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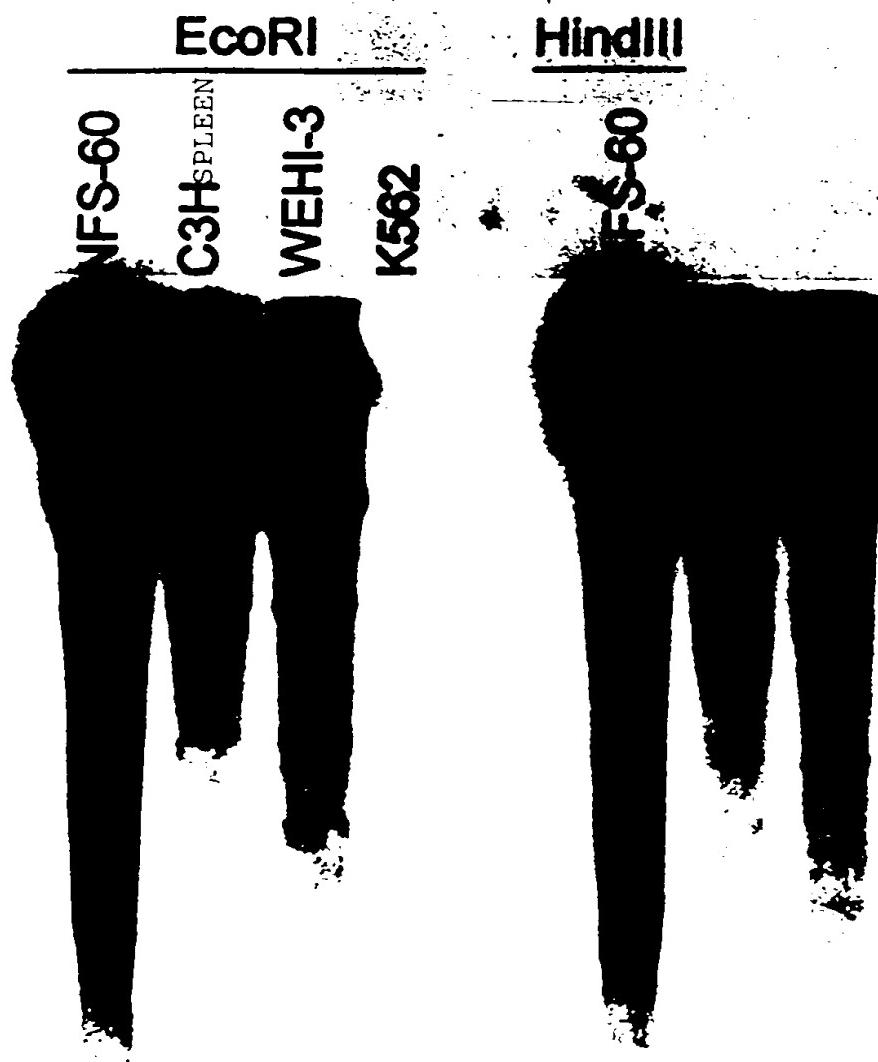
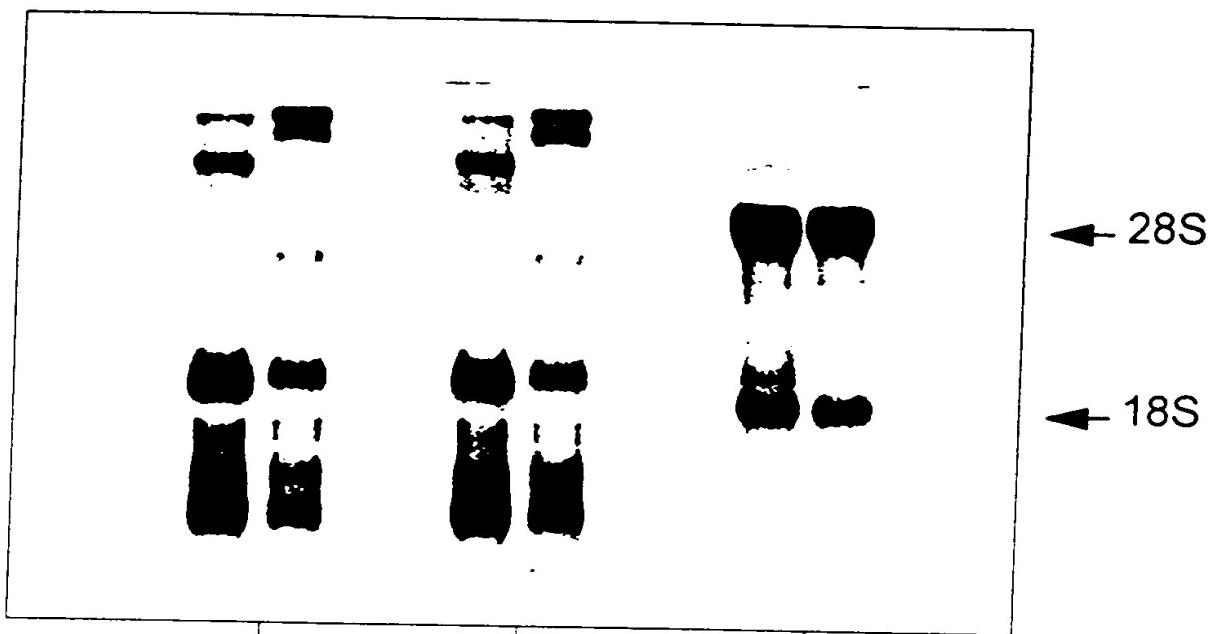


FIGURE 12

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APPROVED BY	O.G. FIG.
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SAMPLE: 1350 bp      500 bp 3'      220 bp 5'

FIGURE 13

APPROVED	O.G. FIG.
BY	CLASS SUBCLASS
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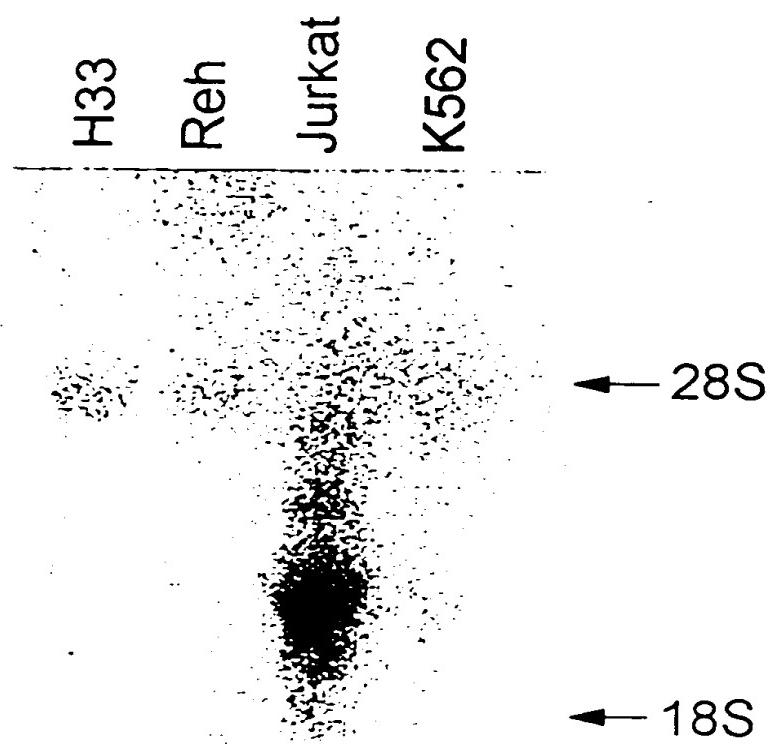


FIGURE 14

APPROVED BY	O.G. FIG.
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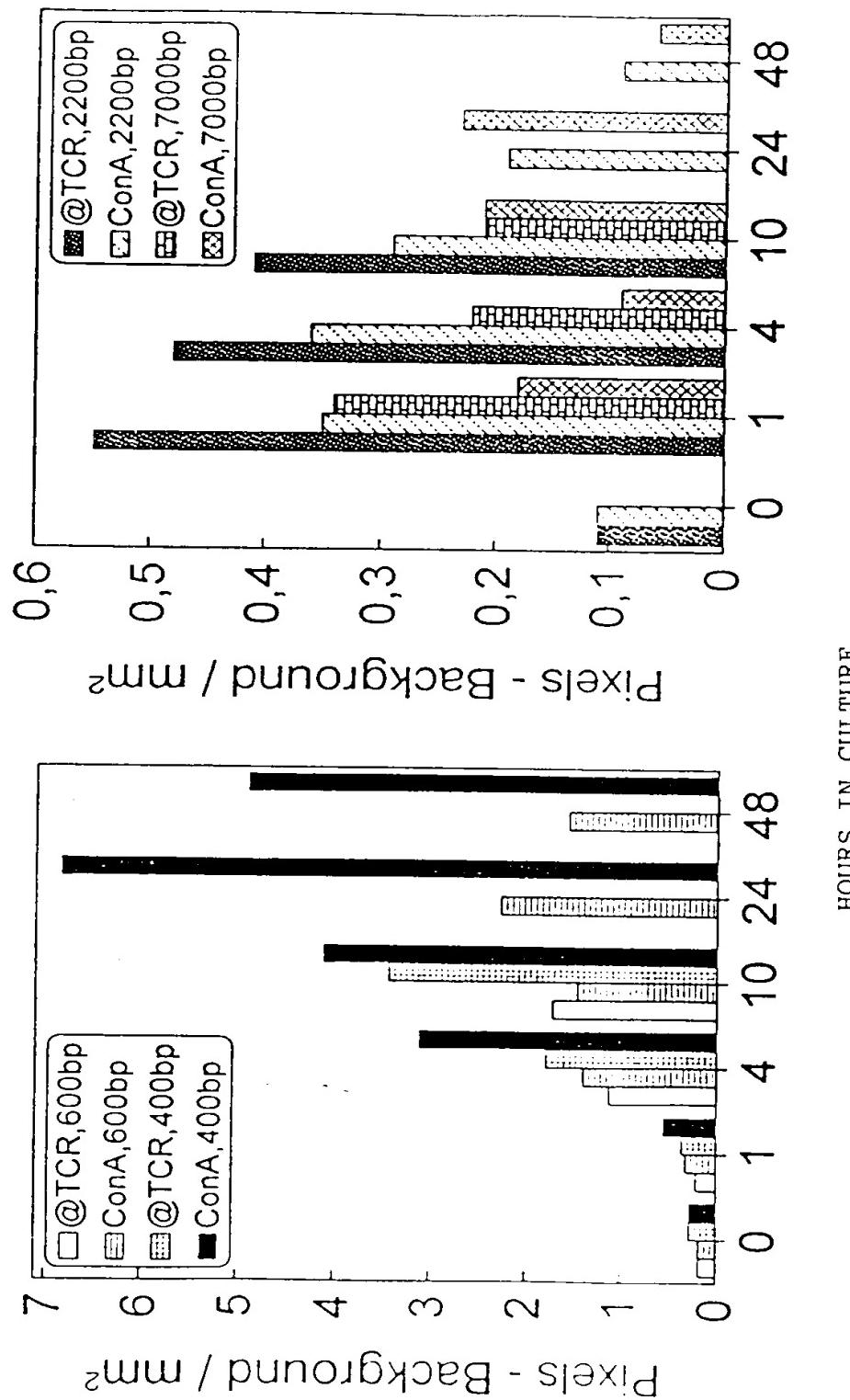


FIGURE 15

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APPROVED	O.G. FIG.
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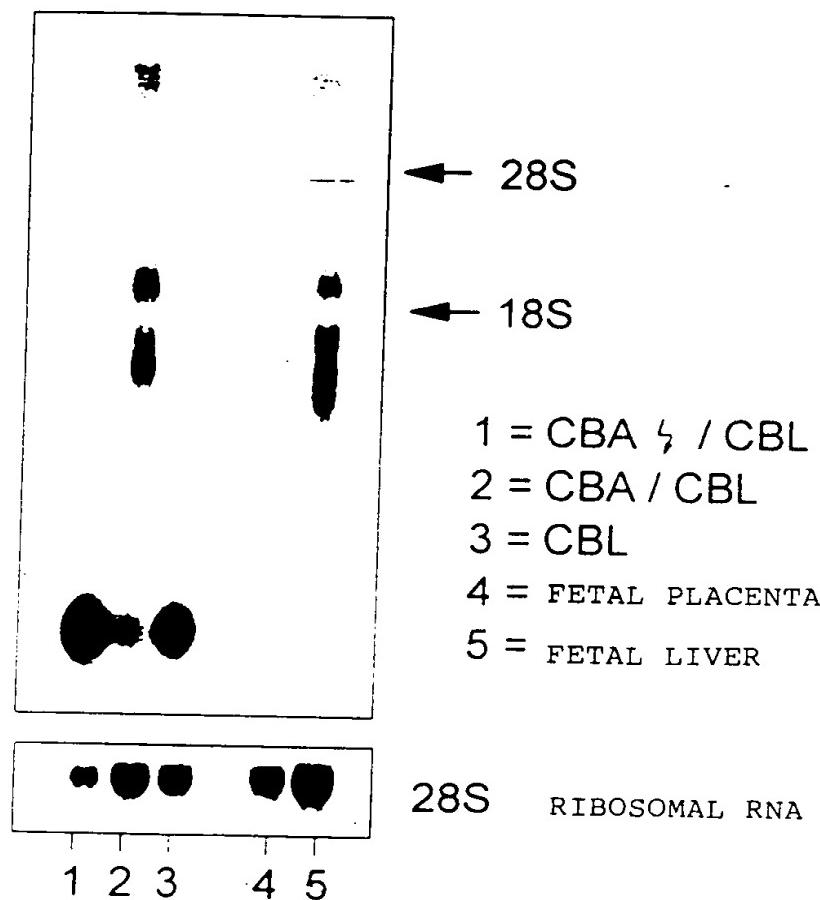


FIGURE 16

APPROVED BY	O.G. FIG.
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TRANSPLANTATION OF CBA X CBL  
OR CBL SPLEEN CELLS

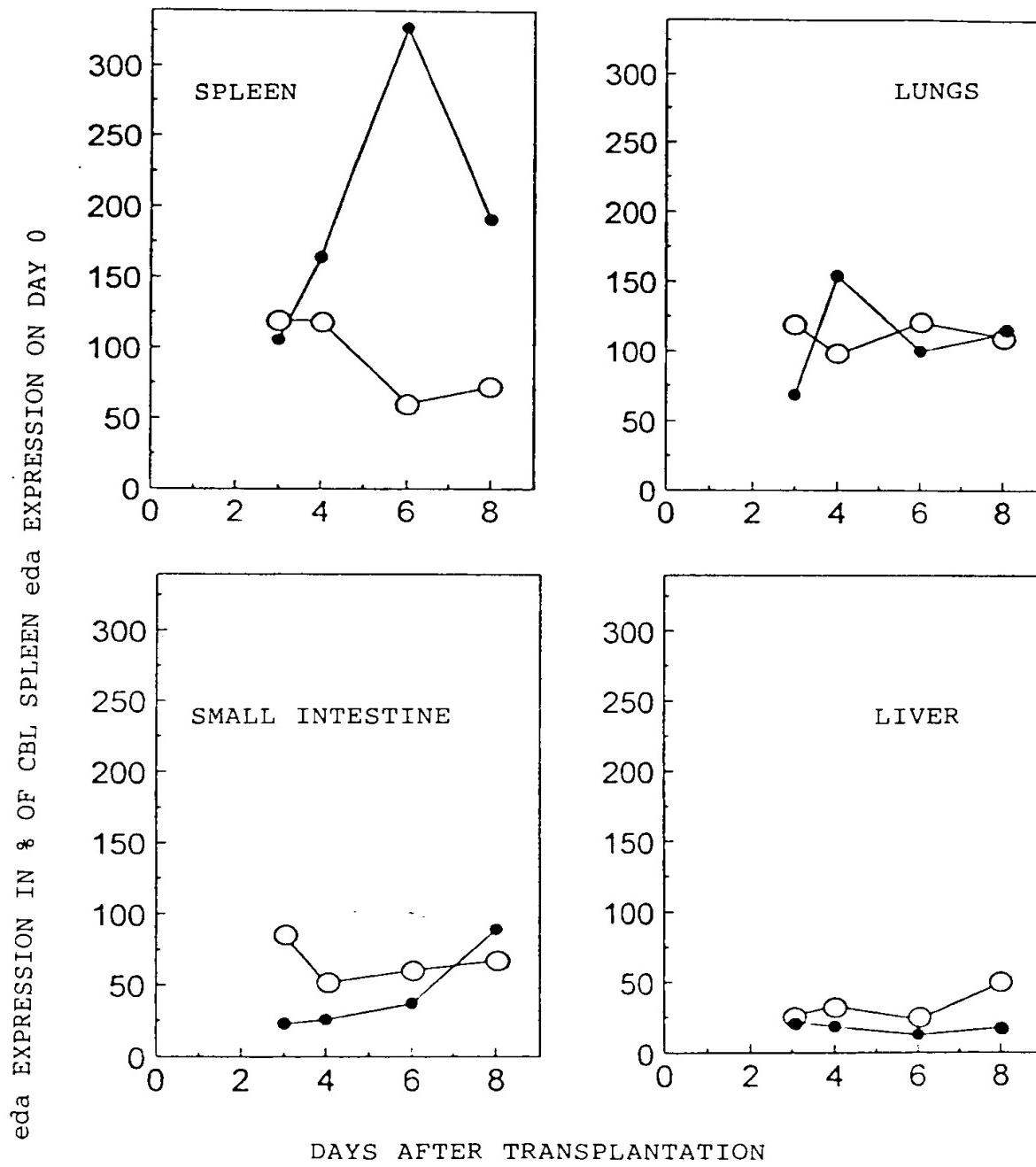


FIGURE 17

APPROVED BY	O.G. FIG.
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1 CCGACCGTGC GGACTTAAGA TGGAGGCAGT TCCTGTCTGC GGCAGGAAGA  
 51 GAAGGCTCGG TCGGAGCCGG GAATGCTGGG ACTTGAGTAGTG CGTAGTCAT  
 101 GGTTCTCTAT GGGCTTCAG AGTAGTGGC GGGAAAGGCGG CCCCAGGGCA  
 151 TGCTGGGAGT TGTAGTCCTG CCGTCGTCAA TGGTTCTCTA TGGGCTTTCA  
 201 GAGTGAGTGG CGGGAAAGGCG GCCCCGAGGC ATGCTGGGAG TTGTAGTCCT  
 251 GCCATAGTCA ATGGTTCTCT ATGGGCTTTC AGACTGAGTG GCAGGAAGGC  
 301 GGCCCCGAGG CATGCTGGGA GTTGCAGCGC CATGTTTAA AGCACCGCGT  
 351 TCTCTGTATA GACCTGGCTG TGGATTTTC GCTAATTCTT TTTTTAGCT  
 401 TTATTTTAA TTTTACTTT TTCACACAGG ATTCTCTTT ATAGCCTTGG  
 451 CTACCGTTT TTCCCTAATT ATTCTCCTT TCATTTGGT TTATTTTTT  
 501 TTAATTTGG TTTTTTAAG ACAGGGTTTC TCTGTATAGA CCTGGCTGTG  
 551 GATTCTCAC TAATTATTT TTTAGCTT ATTAAATT ATTACTTTT  
 601 CACACAGGAT TTCTCTTTAT AGCCTGGCT ACCGTTTTT CCGTAATTAT  
 651 TCTTATTTTC ATTTGGTTT ATTTTTAAT TTTAATT TTTAATT GATTTGGAG  
 701 ACAGGGTTTC TCTTTAGCC GCAGCTATGG TTTCTGCCCT AATTATTCTT  
 751 GTCCTTATTT GTAATTAAAT TCTTAATTAA ATTAAATTAA TAATTTGTT  
 801 GTAAGTTTT CTGTGGCGT GAATGGAAAG TCTAACCGT GTTCTCTGT  
 851 TCAGCGTCCG CCGGTACCGG CCGCCGCC CAGCGACGTC ACCCACACGC  
 901 GCAGAAGCGG ACGCCGCGGT CAAGATGTCT CTGCCATGCC CACGGGACGC  
 951 ACGGACGCAC GGACGGACGG ACGGACTCCA CAAGGTAGGA AGCCTGCGCC  
 1001 GACCGCACCG CCGCACCCAC CACAGCACAC AGGACACACG CGGGCCCCGC  
 1051 GCGCGCCAG GCACACGCGG CACACACGGC ACACACGGCA GGCAGGCCAG  
 1101 GCACACGCAT CCGCAGGACC CGCGCACCC GCCACGCAGA CACGGACGAG  
 1151 CCGCCGCGGT CAAGATGTT ACCCGCCCG GTCAAGATGT ATGTGCCACC  
 1201 GACCCCTCGCC CCGCTGGACG GACGGACGGA CGCACGCACG CGTCAGCGT  
 1251 CCACCGGTCA CTGCCGCCGC CCACAGTGAT GTCACCCACG AAAGCACACA  
 1301 CGTAGAAGCG GACGCCGTGG TCAAGATGTC TCTGCCATCC CCACAGGACG  
 1351 GACGGACGGA CTCCACAAGG TGCCTGTGTC GCCGAGGCCG CCAGGACGGA  
 1401 GCGATTCTCA CGGAGGAAGG AGCACGCCAA CAGGGCCTGA CTGCGTACAG  
 1451 ACATGTCCCC CTCAATAAAA TTGCAGTTGA AATGGAAAAA AAAAAA

FIGURE 18

APPROVED	O.G. FIG.	
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PRELIMINARY STRUCTURE OF eda 2.2 kbp cDNA

\* = stopcodon, R = Repeat

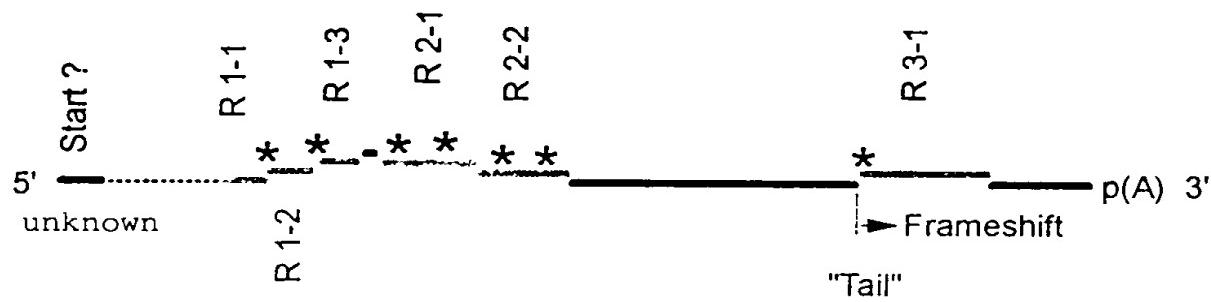


FIGURE 19

APPROVED	O.G. FIG.
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CONSENSUS SEQUENCE

1 CGCGCCCGCC CGGGATCCCC AGCTGCCGCC GCGCCCGCCC GCCCGCCCGG  
51 GGCCCCCGCT GCAGAACCGT GACCGTCCGC CGGTACGGC CGCCGCCCGC  
ATGGGGCT GCAGAACCGT GACCGTCCGC CGGTACGGC CGCCGCCCGC

101 AGCGACGTCA CCCACACGCG CAGAACGGGA CGCCGCCGGTC AAGATGTCTC  
AGCGACGTCA CCCACACGCG CAGAACGGGA CGCCGCCGGTC AAGATGTCTC

151 TGCCATGCC ACAGGGACGCA CGGACGCACG GACGGACGGA CTGACTCCAC  
TGCCATGCC ACAGGGACGCA CGGACGCACG GACGGACGGA CTGACTCCAC

201 AAGGTAGGAA GCCTGCGCCG ACCGCACCGC CGCACCCACC ACAGCACACA  
AAGGTAGGAA GCCTGCGCCG ACCGCACCGC CGCACCCACC ACAGCACACA

251 GGACACACGC GGGCCCCGCG CCCGCCAGG CACACGGCAGC ACACACGGCA  
GGACACACGC GGGCCCCGCG CCCGCCAGG CACACGGCAGC ACACACGGCA

301 CACACGGCAG GCAGGCCAGG CACACGCATC CGCAGGACCC GCCGCACCCG  
CACACGGCAG GCAGGCCAGG CACACGCATC CGCAGGACCC GCCGCACCCG

351 CCACGCAGAC ACGGACGAGC CGCCGCCGGTC AAGATGTTCA CCCGCCGG  
CCACGCAGAC ACGGACGAGC CGCCGCCGGTC AAGATGTTCA CCCGCCGG

401 TCAAGATGTA TGTGCCACCG ACCCTCGCCC CGCTGGACGG ACGGACGGAC  
TCAAGATGTA TGTGCCACCG ACCCTCGCCC CGCTGGACGG ACGGACGGAC

451 GCGGCCACGC CGTCAGCGTC CACCGGTAC TGCCGCCGCC CACAGTGACG  
GCGGCCACGC CGTCAGCGTC CACCGGTAC TGCCGCCGCC CACAGTGACG

501 TCACCCACGA AAGCACACAC GTAGAACGGG ACGCCGTGGT CAAGATGTCT  
TCACCCACGA AAGCACACAC GTAGAACGGG ACGCCGTGGT CAAGATGTCT

551 CTGCCATCCC CACAGGACGG ACGGACGGAC TCCACAAGGT GCGCGTGTG  
CTGCCATCCC CACAGGACGG ACGGACGGAC TCCACAAGGT GCGCGTGTG

601 CCGAGGCCGC CAGGATGGAG CGATTCTCAC GGAGGAAGGA GCACGCCAAC  
CCGAGGCCGC CAGGATGGAG CGATTCTCAC GGAGGAAGGA GCACGCCAAC

651 AGGGCCTGAC TGCACAGA AATGCCCGCC CTCAATAAAA TTGCAGTTGA  
AGGGCCTGAC TGCACAGA AATGCCCGCC CTCAATAAA

701 AATGGAAAAA AAAAAA

APPROVED BY	O.G. FIG.
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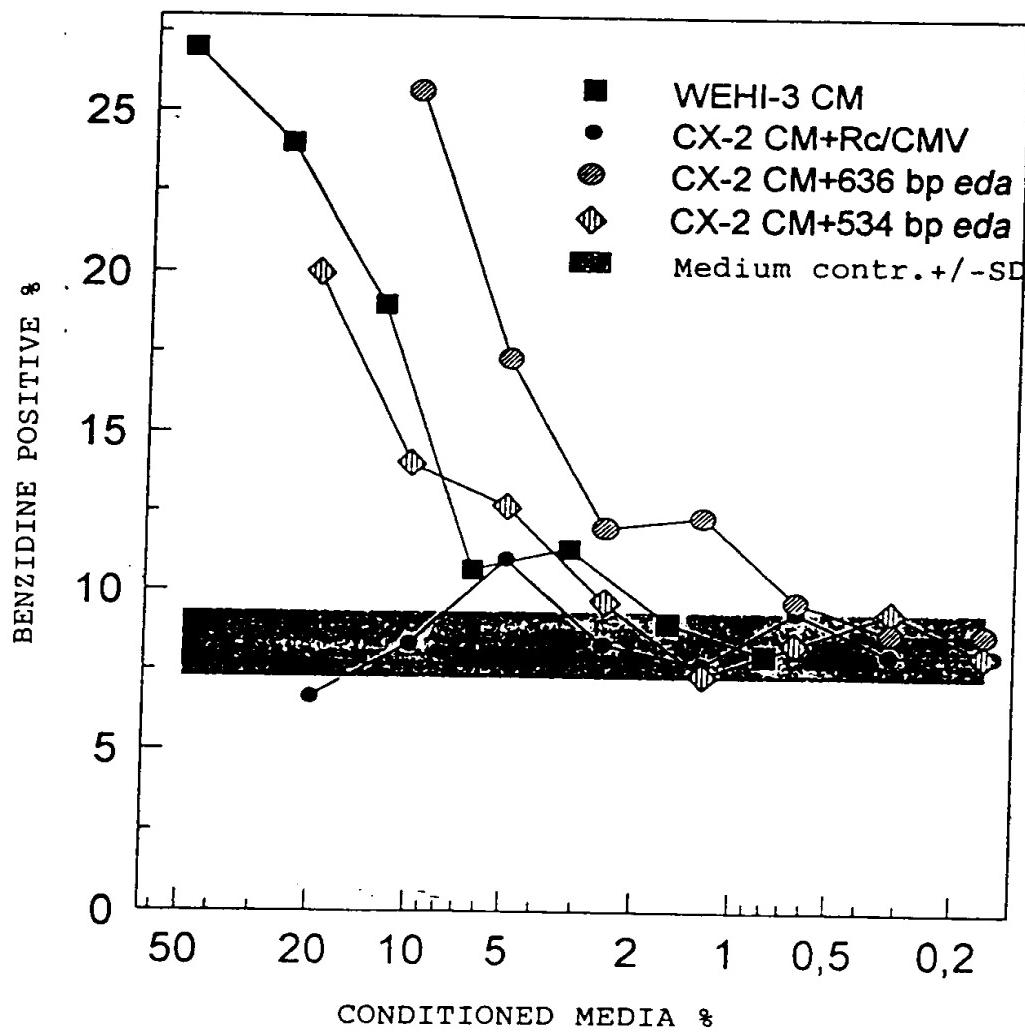
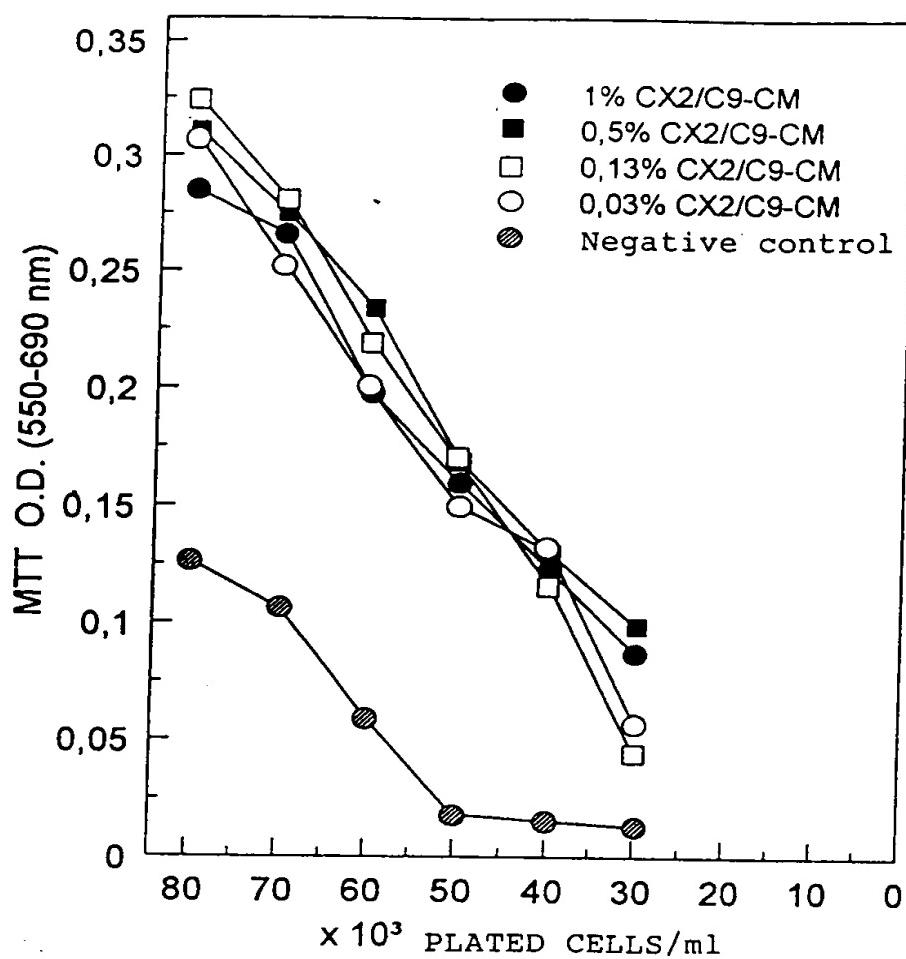


FIGURE 21

APPROVED BY	O.G. FIG.
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A



B

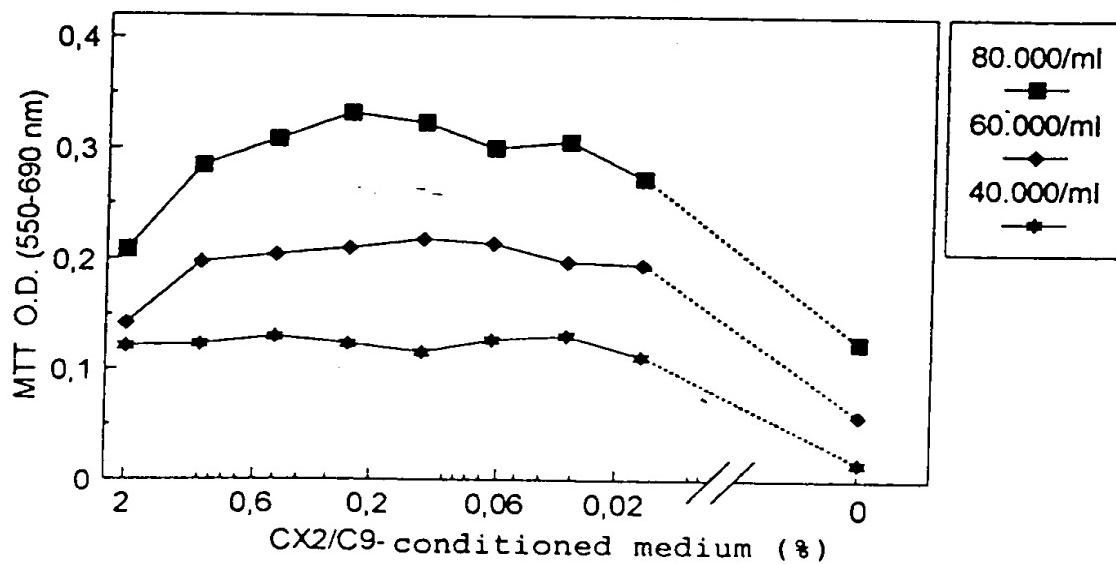


FIGURE 22

APPROVED	O.G. FIG.
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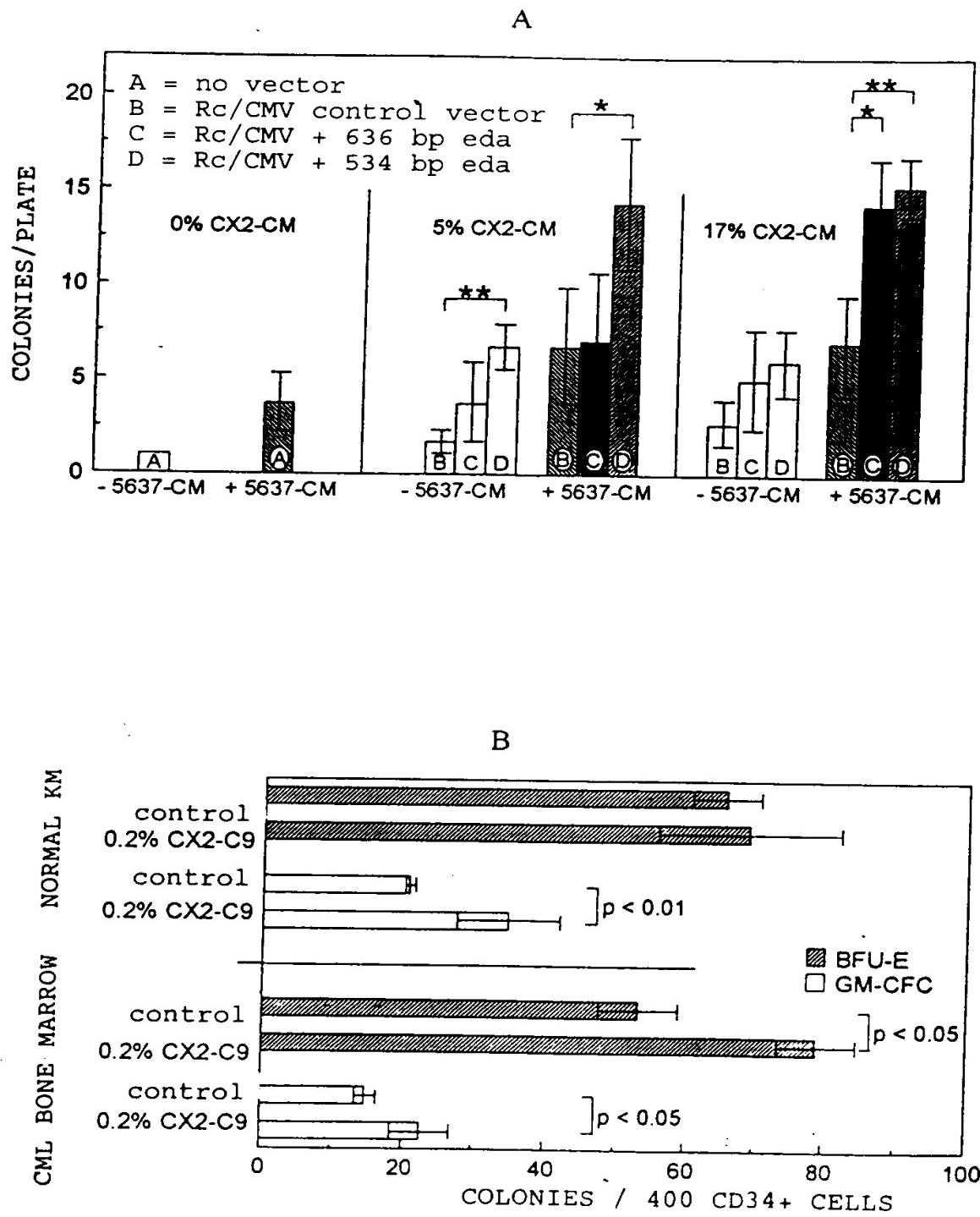


FIGURE 23